PATENT COOPERATION TREATY

PCT

REC'D	2.7	APR	2005
WIPO			PCT

INTERNATIONAL PRELIMINARY REPORT ON PATENTABILITY

(Chapter II of the Patent Cooperation Treaty)

(PCT Article 36 and Rule 70)

				
Applicant's or agent's file reference P036460WO		FOR FURTHER A	CTION	See Form PCT/IPEA/416
International application No. PCT/GB2004/000127		International filing date 14.01.2004	(day/month/year)	Priority date (day/month/year) 14.01.2003
Inter	national Patent Classification (IPC) or	national classification and I	PC	
	2Q1 <i>[</i> 68			
<u></u>				
1	lcant			
TIS	l, Laurence, C. et al.			
1.	Authority under Article 35 and tr	ansmitted to the applicar	nt according to Article	his International Preliminary Examining 36.
2.	This REPORT consists of a total	of 6 sheets, including t	his cover sheet.	
3.	3. This report is also accompanied by ANNEXES, comprising:			
İ	a. 🛭 sent to the applicant and	to the International Bure	au) a total of 4 sheet	ts, as follows:
	sheets of the description, claims and/or drawings which have been amended and are the basis of this report and/or sheets containing rectifications authorized by this Authority (see Rule 70.16 and Section 607 of the Administrative Instructions).			
	beyond the disclosur	ede earlier sheets, but w e in the international app	hich this Authority cor dication as filed, as in	nsiders contain an amendment that goes dicated in item 4 of Box No. I and the
Ì	Supplemental Box.			
	 b. (sent to the International Bureau only) a total of (indicate type and number of electronic carrier(s)), containing a sequence listing and/or tables related thereto, in computer readable form only, as indicated in the Supplemental Box Relating to Sequence Listing (see Section 802 of the Administrative Instructions). 			
				•
4.	This report contains indications	relating to the following i	tems:	
1	☐ Box No. I Basis of the or	pinion		
	☑ Box No. II Priority			
1		ment of opinion with reas	ard to novelty, inventiv	re step and industrial applicability
	☐ Box No. IV Lack of unity of		,,	o otop and manual approachity
Box No. V Reasoned statement under Article 35(2) with regard to novelty, inventive step or industrial applicability; citations and explanations supporting such statement			Ity, inventive step or industrial	
	☐ Box No. VI Certain docum	•	supporting such state	ement
1		s in the international app	lication	
	☐ Box No. VIII Certain observ	• •		
1	Box No. VIII Gertalii observ	auons on the internation	ai application	
Date	of submission of the demand		Date of completion of	this report
15.1	11.2004		26.04.2005	
Nam	e and mailing address of the internation	onai	Authorized Officer	
preiir	minary examining authority: European Patent Office			Charleman and Marie
D-80298 Munich			Hennard, C	
Tel. +49 89 2399 - 0 Tx: 523656 epmu d Fax: +49 89 2399 - 4465			-	2300-7355
			Telephone No. +49 89	7 2000-1000 Pagus . **

INTERNATIONAL PRELIMINARY REPORT ON PATENTABILITY

International application No. PCT/GB2004/000127

	Box	(No. I	Basis of the rep	port			
1.	With filed	With regard to the language , this report is based on the international application in the language in which it wa filed, unless otherwise indicated under this item.					
		This re	port is based on t is the language of	ranslations from the original language into the following language , a translation furnished for the purposes of:			
		☐ pub	lication of the inte	(under Rules 12.3 and 23.1(b)) ernational application (under Rule 12.4) ary examination (under Rules 55.2 and/or 55.3)			
With regard to the elements* of the have been furnished to the receive report as "originally filed" and are			furnished to the r	* of the international application, this report is based on (replacement sheets which eceiving Office in response to an invitation under Article 14 are referred to in this d are not annexed to this report):			
	Des	cription	, Pages				
	1-40			as originally filed .			
	Clai	ms, Nur	nbers				
	1-37			received on 29.03.2005 with letter of 29.03.2005			
	Drav	wings, S	Sheets				
	1/12	-12/12		as originally filed			
	×	a sequ	ence listing and/o	r any related table(s) - see Supplemental Box Relating to Sequence Listing			
3.				resulted in the cancellation of:			
		☐ the	description, page claims, Nos.				
			drawings, sheets, sequence listing				
		□ any	table(s) related to	sequence listing (specify):			
4.	□ had Sup	not bee	port has been est en made, since th tal Box (Rule 70.2	ablished as if (some of) the amendments annexed to this report and listed below ey have been considered to go beyond the disclosure as filed, as indicated in the 2(c)).			
		☐ the	description, page	• ••			
		☐ the	claims, Nos. drawings, sheets				
			sequence listing (table(s) related to	(specify): o sequence listing (specify):			
	*	If ite	em 4 applies,	some or all of these sheets may be marked "superseded."			

INTERNATIONAL PRELIMINARY REPORT ON PATENTABILITY

International application No. PCT/GB2004/000127

	Box No.	II Priority			
1.	preso ⊠ co	cribed time limit the r opy of the earlier app	equested: lication wh	ose priority	ty had been claimed due to the failure to furnish within the has been claimed (Rule 66.7(a)). Deriority has been claimed (Rule 66.7(b)).
2.	been		34.1). Thus	for the pu	ty had been claimed due to the fact that the priority claim has rposes of this report, the international filing date indicated
3.	Additiona	al observations, if nec	essary:		
	Box No.				35(2) with regard to novelty, inventive step or industrial ting such statement
1.	Statemen	nt			
	Novelty (N)	Yes: No:	Claims Claims	1-33, 35 34, 36-37
	Inventive	step (IS)	Yes: No:	Claims Claims	1-33 34-37
	Industrial	applicability (IA)	Yes: No:	Claims Claims	1-37 None
2.	Citations	and explanations (Re	ule 70.7):		
	see sepa	arate sheet			
	Box No.	VII Certain defects	s in the int	ternationa	I application

The following defects in the form or contents of the international application have been noted:

see separate sheet

INTERNATIONAL PRELIMINARY REPORT ON PATENTABILITY

International application No. PCT/GB2004/000127

_								
_	Su	ppl	emental Box relating to Sequence Listing					
C	onti	nua	tion of Box I, item 2:					
1.	 With regard to any nucleotide and/or amino acid sequence disclosed in the international application and necessary to the claimed invention, this report has been established on the basis of: 							
	a. type of material:							
		Ø	a sequence listing					
			table(s) related to the sequence listing					
	b. format of material:							
		Ø	in written format					
		×	in computer readable form					
c. time of filing/furnishing:								
			contained in the international application as filed					
			filed together with the international application in computer readable form					
		×	furnished subsequently to this Authority for the purposes of search and/or examination					
	i	×	received by this Authority as an amendment on					
2.	×	the ad	addition, in the case that more than one version or copy of a sequence listing and/or table(s) relating ereto has been filed or furnished, the required statements that the information in the subsequent or ditional copies is identical to that in the application as filed or does not go beyond the application as filed, appropriate, were furnished.					
3.	. Additional observations, if necessary:							

Re Item V

Reasoned statement under Rule 66.2(a)(ii) with regard to novelty, inventive step or industrial applicability; citations and explanations supporting such statement

1. Reference is made to the following documents:

D1: WO 01/42496 A

D2: ANALYTICAL BIOCHEMISTRY, vol. 288, no. 1, 1 January 2001, pages 28-38

D3: WO 02/064830 A

The comments made by the applicant in his letter dated 15.11.2004 have been taken into consideration for issuing the present IPR.

2. Novelty (Article 33(2) PCT):

D1 (page 14, line 17 - page 16, line 6; page 43, line 30 - page 44, line 33; figures 6-7; claims 1-11) and D2 (page 30, right-hand column; page 32, left-hand column, lines 15-18) disclose a method for determining the amount of nucleic acid in a sample. The method involves mixing a target nucleic acid immobilised on streptavidin-coated beads, extension primers which hybridize to the target, and all the ingredients necessary for amplification of the target (polymerase...), APS, luciferin, ATP sulfurylase and luciferase. The measure of luminescence is made during the amplification using a luminometer. The amount of target nucleic acid is determined using quantitative standards. By modifying claim 1 and specifying that the amplification (step ii) concerns the target nucleic acid and that more than one cycle of amplification are involved, novelty of independent claim 1 is established.

Considering the claimed kit, the documents **D1** and **D2** disclose processes which involve all the components of the kit of **claim 34**. Since from the wording of the claim the kit can be made of only one container (this is further justified by the dependent claim 35 which claims a kit made of two containers) this kit is anticipated by the processes of D1 and D2. Thus **claims 34 and 36** are not novel.

Moreover, the device for performing the method can be represented by a luminometer disclosed in **D1** and **D2** which both disclose a process involving all the elements of the kit of claim 34, therefore **claim 37** is not new.

In the light of these documents, claims 34, 36 and 37 are not new and do not fulfil the requirements of Article 33(2) PCT.

3. Inventive merit (Article 33(3) PCT):

3.1 D1 (same passages as above), which is the closest prior art, concerns a

method for assessing the amount of nucleic acid in a sample involving the measurement of pyrophosphate produced during amplification by means of bioluminescence. The method of **claim 1** of the present application distinguishes itself from the prior art by the amplification which involves more than one amplification cycle.

By performing the process involving more than one amplification cycle, the process of the present **claim 1**, according to the applicant's statement made in his letter dated 15.11.2004, overcomes the prejudice that the elements of the bioluminescence detection are unstable over a longer time period (more than one minute). Therefore, the problem to be solved can be seen in the provision of a method for determining the amount of template nucleic acid present in a sample involving a multiple cycle amplification without loss of bioluminescence activity.

The skilled person would not, knowing that the bioluminescent elements are unstable over time, suggest a method in which multiple amplification cycles are performed. Thus an inventive merit can be recognised for independent claim 1.

- 3.2 Dependent **claim 35**, relating to a kit made of two containers, one containing the ingredients for amplification and the other the ingredients for the bioluminescence is considered not inventive considering the disclosure of **D1** (page 44 and claims 22-23) in combination with **D3** (claim 17).
- 3.5 In order to summarise the above objections, claims 1-33 are considered to involve an inventive merit and to fulfil the requirements of Article 33(3) PCT, whereas claim 35 is not inventive.

4. Industrial applicability (Article 33(4) PCT):

Due to the nature of the claims, an industrial applicability of the invention is obvious and claims 1-37 of the present application are considered to fulfil the requirements of Article 33(4) PCT.

5. Contrary to the requirements of **Rule 5.1(a)(ii) PCT**, the relevant background art disclosed in **D1 and D2** is not mentioned in the description, nor are these documents identified therein.

CLAIMS:

10:22

- 1. A method for determining the amount of template nucleic acid present in a sample comprising the steps of:
- i) bringing into association with the sample all the components necessary for nucleic acid amplification, and all the components necessary for a bioluminescence assay for nucleic acid amplification including:
- a) a nucleic acid polymerase,
- b) the substrates for the nucleic acid polymerase,
- c) at least two primers,
- 10 d) a thermostable luciferase,
 - e) luciferin,

30

- f) optionally ATP sulphurylase, and
- g) optionally adenosine 5' phosphosulphate, and subsequently:
- 15 ii) performing a nucleic acid amplification reaction of the target nucleic acid involving more than one cycle of amplification;
 - iii) monitoring the intensity of light output from the bioluminescence reaction, and
 - iv) determining the amount of template nucleic acid present in the sample.
- 2. A method according to claim 1, wherein at least steps ii) and iii) are carried out in a 20 sealed yessel.
 - 3. A method according to claim 1 or claim 2, wherein in step iii) the intensity of light output is monitored during the nucleic acid amplification reaction.
 - 4. A method according to any one of claims 1 to 3, wherein step iii) further includes producing a data set of intensity of light output as a function of time.
- 25 5. A method according to any one of claims 4 to 6, wherein the amount of template nucleic acid present is determined by measuring from the data set the time taken to reach a point at which the rate of change of intensity of light output changes significantly.
 - 6. A method according to any one of claims 1 to 4 for determining the amount of template nucleic acid present in the sample at the beginning of the nucleic acid amplification reaction of step ii).
 - 7. A method according to any one of claims 1 to 4 for determining the amount of template nucleic acid present in the sample as a result of the nucleic acid amplification reaction of step ii).

10:22

- 8. A method according to any one of claims 5 to 7, wherein the amount of template nucleic acid present is determined by measuring from the data set the time taken to reach a point at which the intensity of light output begins to increase.
- A method according to any one of claims 5 to 7, wherein the amount of template
 nucleic acid present is determined by measuring from the data set the time taken to reach a point at which the intensity of light output is at a maximum.
 - 10. A method according to any one of claims 5 to 7, wherein the amount of template nucleic acid present is determined by measuring from the data set the time taken to reach a point at which the rate of decrease of intensity of light output increases.
- 10 11. A method according to any one of claims 5 to 7, wherein the amount of template nucleic acid present is determined by measuring from the data set the time taken to reach a point at which the rate of decrease of intensity of light output decreases.
- 12. A method according to any one of claims 5 to 7, wherein the amount of template nucleic acid present is determined by measuring from the data set the time taken to reach a point at which the intensity of light output reaches or crosses a predetermined level.
 - 13. A method according to any one of claims 8 to 12, wherein the thermostable luciferase that is brought into association with the sample in step i) is a reversibly-inhibited luciferase.
- 14. A method according to any one of the preceding claims, wherein step iv) further comprises comparing the intensity of light output to the intensity of light output from a control in which the sample comprises a known amount of template nucleic acid.
 - 15. A method according to any one of claims 1 to 14 for determining whether the template nucleic acid is present in the sample.
- 16. A method according to claim 14, wherein whether the template nucleic acid is present in the sample is determined by measuring from the data set whether the intensity of light output reaches or crosses a predetermined level.
 - 17. A method according to claim 15, wherein an increase in the intensity of light output relative to the predetermined level indicates the presence of template nucleic acid in the sample.
- 30 18. A method according to claim 15, wherein a decrease in the intensity of light output relative to the predetermined level indicates the presence of template nucleic acid in the sample.

20

10:22

- 19. A method according to any one of claims 16 to 18, wherein whether the template nucleic acid is present in the sample is determined by measuring from the data set whether the intensity of light output reaches or crosses the predetermined level within a predetermined length of time following the start of the amplification reaction of step ii).
- 5 20. A method according to any one of the preceding claims, wherein step iv) further comprises comparing the intensity of light output to the intensity of light output from a control in which no amplification has taken place.
- 21. A method according to claim 20, wherein a decrease in the intensity of light output relative to a control reaction in which no amplification has taken place indicates the presence of template nucleic acid in the sample.
 - 22. A method according to any one of claims 1 to 21, wherein the nucleic acid amplification reaction of step ii) is a low temperature thermocycling amplification method in which the cycling temperature range does not exceed 75°C.
- 23. A method according to any one of claims 1 to 21, wherein the nucleic acid amplification reaction of step ii) is carried out isothermally.
 - 24. A method according to claim 23, wherein the nucleic acid amplification reaction of step ii) is carried out within a temperature range that does not exceed 75°C.
 - 25. A method according to claim 23 or claim 24, wherein the nucleic acid amplification reaction of step ii) is carried out at a constant temperature at which the components of the amplification reaction and the bioluminescence assay are stable.
 - 26. A method according to claim 23 or claim 24, wherein the nucleic acid amplification reaction of step ii) is carried out at more than one temperature within the temperature range in which the components of the amplification reaction and the bioluminescence assay are stable.
- 25 27. A method according to claim 26, wherein the nucleic acid amplification reaction of step ii) is started at a higher temperature and subsequently dropped to a lower temperature.
 - A method according to any preceding claim for use in medical diagnostics.
 - 29. A method according to any preceding claim for use in determining whether a pathogen is present in a sample.
- 30 30. A method according to any preceding claim for determining whether a particular nucleic acid sequence is present in an organism's genetic code.
 - 31. A method according to claim 30 for determining whether the nucleic acid to which the template nucleic acid originates has been genetically modified.

10:22

44

- 32. A method according to any one of claims 1 to 27 for determining whether an organism is present in a sample.
- 33. A method according to any one of claims 1 to 27 for use in immuno-nucleic acid amplification technology.
- A kit for use in a method according to any one of the preceding claims, wherein the kit comprises a nucleic acid polymerase, the substrates for the nucleic acid polymerase, at least two primers, a thermostable luciferase, luciferin and optionally ATP sulphurylase and adenosine 5' phosphosulphate.
- 35. A kit for use in a method according to any one of claims 1 to 33, wherein the kit comprises containers respectively containing:
 - a) a buffered mixture of nucleic acid polymerase, a source of Mg and dNTPs; and b) a luciferase, luciferin and ATP sulphurylase.
 - 36. A kit according to claim 34 or claim 35, wherein at least one of the components of the kit is in a form which is suitable for storage in the kit.
- 15 37. A device for performing a method according to any one of claims 1 to 32, wherein said device incorporates the components that are present in a kit according to any one of claims 34 to 36.